

SEQUENCE LISTING

#110: Labow, Mark A.
Mickanin, Craig Stephen
Bhatia, Umesh

#120: MAMMARY GLAND CHEMOKINE

#130: 12345

#160: 1

#170: FastSEQ for Windows Version 4.0

#210: 1

#211: 1017

#212: DNA

#213: HUMAN

#400: 1

tagataccct	gaacacotcc	cagggcgggg	ccacctgget	tacttttcc	ctgcacttcc	60
tctgtgcaca	aggacacott	tagcctcatt	tcctgatcga	acagctccac	ttgtgttgc	120
gtcagtgcac	gtagggcagg	caggaatgca	gcagagagga	ctggccatcg	tggtcttggc	180
tgtctgtggg	gocctacatg	cctcagaagg	catacttccc	attgctccca	gctgttgac	240
ggaggtttca	catacatatt	ccagaaggct	cctggaaaga	gtgaatatgt	gtgcaccca	300
gagagctgat	ggggattgtg	acttggctgc	tgtcatctct	cattgcacgc	gcagaagaat	360
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ataaatctac	agagacaatt	cctcaagtgg	acttggccat	gattggttgt	aagtttatca	600
tctgaattct	ccttattgta	gacaacagaa	caaaaacaaaa	tattggtttt	taaaaaatga	660
acaattgtgc	gggtatgcaaa	tgtagccaat	aataatactca	aactcctggg	ctcaagcgat	720
cctcccaact	tagctctcca	aagtaactggg	attataggtg	tgagccacag	tgcttggcct	780
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aatagtttta	actaaatata	acttcaaaaac	gtctagtgtg	agtagctaac	gttgtttgga	960
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ccgctgattt	ttgtattttc	agtagagacg	gggttttccc	acgttggccg	ggctggctcc	1380
aaactcttga	cctcaagtga	accacccgcb	tgtgcctccc	aaagtgtctg	aattaccagc	1440
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agaaagcaat	agagagccag	agccacttta	tcaggtggca	ggtgtcccg	gcctccctgc	1620
tggttagtcc	caagcgggtg	tgttgccagg	atgtcttggc	ggtgataatg	ggacacacag	1680

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aggcaatgag totocatagg ttaaaatgcc accaaaaactg gcttttgct aatatccctc 1740
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tatttccact ttatagatga ggaaatttga ggcctcttaga gytaaaatga cttgcccagg 1860
ccacacagga agtggcagag acaagctttt taaataagaa aaaattaata aaatataata 1920
ccagagtaac ttaaaatatt aataaaccac aattttaaat taattaaccg tgataaccac 1980
cattaataaa agttaagata ccaaaaaaaaa aaaaaaa 2017

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4010: 2
 4011: 127
 4012: PRT
 4013: HUMAN

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4900: 2
Met Gln Gln Arg Gly Leu Ala Ile Val Ala Leu Ala Val Cys Ala Ala
1 5 10 15
Leu His Ala Ser Glu Ala Ile Leu Pro Ile Ala Ser Ser Cys Cys Thr
20 25 30
Glu Val Ser His His Ile Ser Arg Arg Leu Leu Glu Arg Val Asn Met
35 40 45
Cys Arg Ile Gln Arg Ala Asp Gly Asp Cys Asp Leu Ala Ala Val Ile
50 55 60
Leu His Val Lys Arg Arg Arg Ile Cys Val Ser Pro His Asn His Thr
65 70 75 80
Val Lys Gln Trp Met Lys Val Gln Ala Ala Lys Lys Asn Gly Lys Gly
85 90 95
Asn Val Cys His Arg Lys Lys His His Gly Lys Arg Asn Ser Asn Arg
100 105 110
Ala His Gln Gly Lys His Glu Thr Tyr Gly His Lys Thr Pro Tyr
115 120 125

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